

CLAIM AMENDMENTS

Claims 1 through 90 (canceled)

1 91. (New) An isolated polynucleotide coding for a
2 polypeptide comprising the amino acid sequence of SEQ ID NO:2.

1 92. (New) A vector comprising the isolated polynucleotide
2 of claim 91.

1 93. (New) A bacterium of the genus *Corynebacterium*
2 comprising the isolated polynucleotide of claim 91.

1 94. (New) The bacterium of claim 93, wherein said
2 bacterium is one of the species *Corynebacterium glutamicum*.

1 95. (New) A bacterium of the species *Escherichia coli* or
2 *Corynebacterium glutamicum* comprising the vector of claim 92.

1 96. (New) An isolated polynucleotide comprising the
2 nucleotide sequence of nucleotides 165 to 3587 of SEQ ID NO: 1.

1 97. (New) A vector comprising the isolated polynucleotide
2 of claim 96.

1 98. (New) A bacterium of the genus *Corynebacterium*
2 comprising the isolated polynucleotide of claim 96.

1 99. (New) The bacterium of claim 98, wherein said
2 bacterium is one of the species *Corynebacterium glutamicum*.

1 100. (New) A bacterium of the species *Escherichia coli* or
2 *Corynebacterium glutamicum* comprising the vector of claim 97.

1 101. (New) An isolated polynucleotide comprising the
2 nucleotide sequence of SEQ ID NO:1.

1 102. (New) A vector comprising the isolated
2 polynucleotide of claim 101.

1 103. (New) A bacterium of the genus *Corynebacterium*
2 comprising the isolated polynucleotide of claim 101.

1 104. (New) The bacterium of claim 103, wherein said
2 bacterium is one of the species *Corynebacterium glutamicum*.

1 105. (New) A bacterium of the species *Escherichia coli* or
2 *Corynebacterium glutamicum* comprising the vector of claim 102.

1 106. (New) An isolated polypeptide having pyruvate
2 carboxylase enzymatic activity comprising the amino acid sequence
3 of SEQ ID NO: 2.

1 107. (New) A pVWEX1pyc vector deposited under DSM 12893.

1 108. (New) A bacterium comprising the vector of claim
2 107.

1 109. (New) An isolated pyruvate carboxylase polypeptide
2 having an amino acid sequence at least 95% identical to the amino
3 acid sequence of the pyruvate carboxylase polypeptide having the
4 complete amino acid sequence in SEQ ID NO:2.